

**AMENDMENTS TO THE SPECIFICATION:**

Please amend the specification as follows:

Please amend the paragraph beginning at line 15 of page 9 of the specification as follows:

For each primary transformant which has proved to be in accordance with the expected phenotype expected from the introduced transgene and which has been selected according to the criteria - monolocus or monocopy and absence of extraborders - the genomic sequences of the host adjacent to the T-DNA may be isolated and identified, for example via a method based on PCR (Polymerase Chain Reaction, Saiki Rk RK. et al., 1988), preferably IPCR (Inverse PCR, Does Mp. Et MP. et al., 1991); the aim being to identify the parental origin of the genome which has accepted the transgene (line of agronomic interest or transformation line).

Please amend the subheading at page 26 of the specification as follows:

**LEGENDS TO THE FIGURES BRIEF DESCRIPTION OF THE DRAWINGS**

Please amend the paragraph beginning at line 25 of page 47 of the specification as follows:

The result of the hybridization is represented on Figure 3: on this autoradiograph indicates that, on an autoradiograph, DNA corresponding to 7 different transformants is visualized, in the knowledge that there are from 1 to 3 plants per transformant.